

Julian R Dupuis

List of Publications by Year in descending order

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Version: 2024-02-01

33
papers

1,199
citations

687220

13
h-index

434063

31
g-index

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all docs

36
docs citations

36
times ranked

2622
citing authors

#	ARTICLE	IF	CITATIONS
1	Gauging ages of tiger swallowtail butterflies using alternate SNP analyses. <i>Molecular Phylogenetics and Evolution</i> , 2022, 171, 107465.	1.2	2
2	Genomic Data Support the Elevation of the Federally Listed El Segundo Blue (<i>Euphilotes</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 702 Td (l	0.6	1
3	Genomic-wide sequencing reveals remarkable connection between widely disjunct populations of the internationally threatened bog buck moth. <i>Insect Conservation and Diversity</i> , 2020, 13, 495-500.	1.4	4
4	Phylogenomic test of mitochondrial clues to archaic ancestors in a group of hybridizing swallowtail butterflies. <i>Molecular Phylogenetics and Evolution</i> , 2020, 152, 106921.	1.2	7
5	Gene flow and climate-associated genetic variation in a vagile habitat specialist. <i>Molecular Ecology</i> , 2020, 29, 3889-3906.	2.0	19
6	Disjunction between canola distribution and the genetic structure of its recently described pest, the canola flower midge (<i>Contarinia brassicola</i>). <i>Ecology and Evolution</i> , 2020, 10, 13284-13296.	0.8	5
7	Does hunger lead to hybridization in a genus of sexually cannibalistic insects (Orthoptera:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50	0.7	1
8	Confidently identifying the correct K value using the $\hat{\pi}$ K method: When does $K = \hat{A}^2$?. <i>Molecular Ecology</i> , 2020, 29, 862-869.	2.0	67
9	Genomics confirms surprising ecological divergence and isolation in an endangered butterfly. <i>Biodiversity and Conservation</i> , 2020, 29, 1897-1921.	1.2	11
10	Population genomic and phenotype diversity of invasive <i>Drosophila suzukii</i> in Hawai'i. <i>Biological Invasions</i> , 2020, 22, 1753-1770.	1.2	14
11	Phylogenomics reveals conservation challenges and opportunities for cryptic endangered species in a rapidly disappearing desert ecosystem. <i>Biodiversity and Conservation</i> , 2020, 29, 2185-2200.	1.2	3
12	Range-wide population genomics of the Mexican fruit fly: Toward development of pathway analysis tools. <i>Evolutionary Applications</i> , 2019, 12, 1641-1660.	1.5	12
13	Targeted amplicon sequencing of 40 nuclear genes supports a single introduction and rapid radiation of Hawaiian <i>Metrosideros</i> (Myrtaceae). <i>Plant Systematics and Evolution</i> , 2019, 305, 961-974.	0.3	15
14	Environmental effects on gene flow in a species complex of vagile, hilltopping butterflies. <i>Biological Journal of the Linnean Society</i> , 2019, 127, 417-428.	0.7	6
15	A new species of <i>Contarinia</i> Rondani (Diptera: Cecidomyiidae) that induces flower galls on canola (Brassicaceae) in the Canadian prairies. <i>Canadian Entomologist</i> , 2019, 151, 131-148.	0.4	8
16	Mitochondrial phylogenomics, the origin of swallowtail butterflies, and the impact of the number of clocks in Bayesian molecular dating. <i>Systematic Entomology</i> , 2018, 43, 460-480.	1.7	34
17	HiMAP: Robust phylogenomics from highly multiplexed amplicon sequencing. <i>Molecular Ecology Resources</i> , 2018, 18, 1000-1019.	2.2	30
18	Genomics-informed species delimitation to support morphological identification of anglewing butterflies (Lepidoptera: Nymphalidae: Polytonia). <i>Zoological Journal of the Linnean Society</i> , 2018, 183, 372-389.	1.0	1

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19	<sc>mvmapper</sc>: Interactive spatial mapping of genetic structures. Molecular Ecology Resources, 2018, 18, 362-367.	2.2	2
20	Population genomics and comparisons of selective signatures in two invasions of melon fly, <i>Bactrocera cucurbitae</i> (Diptera: Tephritidae). Biological Invasions, 2018, 20, 1211-1228.	1.2	19
21	Phylogenomics supports incongruence between ecological specialization and taxonomy in a charismatic clade of buck moths. Molecular Ecology, 2018, 27, 4417-4429.	2.0	13
22	Molecular Characterization of the 2016 New World Screwworm (Diptera: Calliphoridae) Outbreak in the Florida Keys. Journal of Medical Entomology, 2018, 55, 938-946.	0.9	7
23	Would an <sc>RRS</sc> by any other name sound as <sc>RAD</sc>?. Methods in Ecology and Evolution, 2018, 9, 1920-1927.	2.2	27
24	Genomic data indicate ubiquitous evolutionary distinctiveness among populations of California metalmark butterflies. Conservation Genetics, 2018, 19, 1097-1108.	0.8	8
25	The $K = 2$ conundrum. Molecular Ecology, 2017, 26, 3594-3602.	2.0	454
26	Genome-wide SNPs resolve phylogenetic relationships in the North American spruce budworm (<i>Choristoneura fumiferana</i>) species complex. Molecular Phylogenetics and Evolution, 2017, 111, 158-168.	1.2	32
27	Cross-platform compatibility of <i>de novo</i> -aligned <sc>SNP</sc>s in a nonmodel butterfly genus. Molecular Ecology Resources, 2017, 17, e84-e93.	2.2	14
28	The latitudinal diversity gradient in New World swallowtail butterflies is caused by contrasting patterns of out-of and into the tropics dispersal. Global Ecology and Biogeography, 2017, 26, 1447-1458.	2.7	24
29	Hybrid dynamics in a species group of swallowtail butterflies. Journal of Evolutionary Biology, 2016, 29, 1932-1951.	0.8	13
30	Repeated Reticulate Evolution in North American <i>Papilio machaon</i> Group Swallowtail Butterflies. PLoS ONE, 2015, 10, e0141882.	1.1	25
31	Genetic evaluation of the evolutionary distinctness of a federally endangered butterfly, <i>Lange's Metalmark</i> . BMC Evolutionary Biology, 2015, 15, 73.	3.2	14
32	Multi-locus species delimitation in closely related animals and fungi: one marker is not enough. Molecular Ecology, 2012, 21, 4422-4436.	2.0	269
33	Trogus parasitoids of <i>Papilio</i> butterflies undergo extended diapause in western Canada (Hymenoptera.) Tj ETQq1 1 0,784314,rgBT /O	0.8	26